

SEQUENCE LISTING

(1) GENERAL INFORMATION: (i) APPLICANT: Yechezkel Kashi et al. (ii) TITLE OF INVENTION: ABUNDANT, WELL DISTRIBUTED AND HYPERPOLYMORPHIC SIMPLE SEQUENCE REPERIOR OF SAME PROKARYOTE GENOMES AND USE OF SAME PROKARYOTE CLASSIFICATION AND TYPING	.TS
(ii) TITLE OF INVENTION: ABUNDANT, WELL DISTRIBUTED AND HYPERPOLYMORPHIC SIMPLE SEQUENCE REPEI IN PROKARYOTE GENOMES AND USE OF SAME PROKARYOTE CLASSIFICATION AND TYPING	.TS
HYPERPOLYMORPHIC SIMPLE SEQUENCE REPEI IN PROKARYOTE GENOMES AND USE OF SAME PROKARYOTE CLASSIFICATION AND TYPING	TS
IN PROKARYOTE GENOMES AND USE OF SAME PROKARYOTE CLASSIFICATION AND TYPING	113
PROKARYOTE CLASSIFICATION AND TYPING	
	FOR
(iii) NUMBER OF SEQUENCES: 42	
(iv) CORRESPONDENCE ADDRESS:	
(A) ADDRESSEE: Mark M. Friedman c/o Anthony Castorina	
(B) STREET: 2001 Jefferson Davis Highway, Suite 207	
(C) CITY: Arlington	
(D) STATE: Virginia	
(E) COUNTRY: United States of America	
(F) ZIP: 22202	
(v) COMPUTER READABLE FORM:	
(A) MEDIUM TYPE: 1.44 megabyte, 3.5" microdisk	
(B) COMPUTER: Twinhead® Slimnote-890TX	
(C) OPERATING SYSTEM: MS DOS version 6.2,	
Windows version 3.11	
(D) SOFTWARE: Word for Windows version 2.0 converted	.0
an ASCI file	
(vi) CURRENT APPLICATION DATA:	
(A) APPLICATION NUMBER:	
(B) FILING DATE:	
(C) CLASSIFICATION:	
(vii) PRIOR APPLICATION DATA:	
(A) APPLICATION NUMBER:	
(B) FILING DATE:	
(viii) ATTORNEY/AGENT INFORMATION: (A) NAME Friedmam, Mark M.	
(6)	
(C) ALL EXECUTE TO SERVICE TO SER	
072.2.562552	
(A) TELEPHONE: 972-3-5625553 (B) TELEFAX: 972-3-5625554	
(C) TELEX:	
(C) IEDDA.	
(2) INFORMATION FOR SEQ ID NO:1:	
(i) SEQUENCE CHARACTERISTICS:	
(A) LENGTH: 23	
(B) TYPE: nucleic acid	
(C) STRANDEDNESS: single	
(D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:	
GATTTTGCAT ATGAGTATAT TAC 23	
عد	
(2) INFORMATION FOR SEQ ID NO:2:	
(i) SEQUENCE CHARACTERISTICS:	
(A) LENGTH: 21	
(B) TYPE: nucleic acid	
(C) STRANDEDNESS: single	
(D) TOPOLOGY: linear	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:	
TTAATTACAG GATGTTCAGT C 21	

- (2) INFORMATION FOR SEQ ID NO:2:
 - (i) SEQUENCE CHARACTERISTICS:





```
LENGTH:
                                      nucleic acid
                       TYPE:
               (B)
                       STRANDEDNESS:
                                      single
               (C)
                                      linear
                       TOPOLOGY:
               (D)
               SEQUENCE DESCRIPTION: SEQ ID NO:3:
       (xi)
AATTTATCCG GTGAATGTGG T 21
```

INFORMATION FOR SEQ ID NO:4: (2)

SEQUENCE CHARACTERISTICS: (i)

LENGTH: (A)

TYPE: (B)

nucleic acid

STRANDEDNESS: single (C) TOPOLOGY:

linear

(D) SEQUENCE DESCRIPTION: SEQ ID NO:4: (xi)

CAACTTAATC TCGGGCTGAC 20

INFORMATION FOR SEQ ID NO:5: (2)

SEQUENCE CHARACTERISTICS:

LENGTH: (A)

TYPE: (B)

nucleic acid

STRANDEDNESS: single (C)

TOPOLOGY:

linear

(D) SEQUENCE DESCRIPTION: SEQ ID NO:5:

(xi) TACATGGCTG ATTATGCGG 20

INFORMATION FOR SEQ ID NO:6: (2)

SEQUENCE CHARACTERISTICS: (i)

> LENGTH: (A)

TYPE:

nucleic acid

(C)

STRANDEDNESS: single

TOPOLOGY: (D)

linear

SEQUENCE DESCRIPTION: SEQ ID NO:6: (xi)

TCGCTATGAA TATCTACTGA C 21

INFORMATION FOR SEQ ID NO:7: (2)

SEQUENCE CHARACTERISTICS: (i)

TYPE:

nucleic acid

(C)

STRANDEDNESS: single TOPOLOGY:

LENGTH:

(D) SEQUENCE DESCRIPTION: SEQ ID NO:7:

GTCAGAGCAG ATCCAGAATG 20

INFORMATION FOR SEQ ID NO:8: (2)

SEQUENCE CHARACTERISTICS:

(A)

LENGTH:

TYPE: (B)

nucleic acid

STRANDEDNESS: single (C)

TOPOLOGY: (D)

linear

SEQUENCE DESCRIPTION: SEQ ID NO:8: (xi)

TCTACAGCAA ATGAACAATG 20

INFORMATION FOR SEQ ID NO:9: (2)

SEQUENCE CHARACTERISTICS: (i)

> (A) LENGTH:

20

(B)

nucleic acid

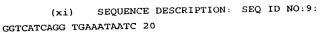
STRANDEDNESS: (C)

single

TOPOLOGY: (D)

linear





INFORMATION FOR SEQ ID NO:10: (2)

- SEQUENCE CHARACTERISTICS: (i)
 - LENGTH: (A)

TYPE: (B)

nucleic acid

STRANDEDNESS: single (C)

(D) TOPOLOGY: linear

SEQUENCE DESCRIPTION: SEQ ID NO:10: (xi)

CGTCCTGATA GATAAAGTGT C 21

INFORMATION FOR SEQ ID NO:11: (2)

- SEQUENCE CHARACTERISTICS:
 - LENGTH:

TYPE: (B)

nucleic acid

STRANDEDNESS: single (C)

TOPOLOGY: (D)

linear

SEQUENCE DESCRIPTION: SEQ ID NO:11: (xi)

CAATGGAACT TACCAATGAC 20

- INFORMATION FOR SEQ ID NO:12: (2)
 - SEQUENCE CHARACTERISTICS:

LENGTH: (A)

20

TYPE: (B)

nucleic acid

STRANDEDNESS: single (C)

TOPOLOGY: (D)

linear SEQUENCE DESCRIPTION: SEQ ID NO:12:

TACCGCGAAG AATTCAACAC 20

(xi)

INFORMATION FOR SEQ ID NO:13: (2)

SEQUENCE CHARACTERISTICS:

LENGTH: (A)

TYPE: (B)

nucleic acid

(C)

STRANDEDNESS: single

(D)

TOPOLOGY: linear

(xi)

SEQUENCE DESCRIPTION: SEQ ID NO:13:

AGCATCAGCG CACAATGCAC 20

- INFORMATION FOR SEQ ID NO:14: (2)
 - SEQUENCE CHARACTERISTICS: (i)

LENGTH:

(A)

TYPE:

nucleic acid

(C)

STRANDEDNESS: single

TOPOLOGY: (D)

linear

SEQUENCE DESCRIPTION: SEQ ID NO:14: (xi)

TGTATGCAGG_CTGGCACAAC 20

- INFORMATION FOR SEQ ID NO:15: (2)
 - SEQUENCE CHARACTERISTICS: LENGTH: (A)

(B)

nucleic acid

STRANDEDNESS: single (C)

linear

TOPOLOGY: (D)

SEQUENCE DESCRIPTION: SEQ ID NO:15: (xi)

ATAACGATCT CCATATCTAC 20

INFORMATION FOR SEQ ID NO:16: (2)





(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH:

20

(B) TYPE:

nucleic acid

(C) STRANDEDNESS: single

(D) TOPOLOGY:

linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:16:

CTCTATCAGC AACTTCTGCC 20

(2) INFORMATION FOR SEQ ID NO:17:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH:

20

(B) TYPE:

nucleic acid

(C) STR

STRANDEDNESS: single

(D) TOPOLOGY:

linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:17:

ATCCGCAGGA TTTTCGCACC 20

(2) INFORMATION FOR SEQ ID NO:18:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH:

20

(B) TYPE:

nucleic acid

(C) ST

STRANDEDNESS: single

(D) TOPOLOGY:

linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:18:

TGCCAGCGTA AATCCGCAAC 20

(2) INFORMATION FOR SEQ ID NO:19:

(i) SEQUENCE CHARACTERISTICS:

(A) (B)

LENGTH:

TYPE:

2

nucleic acid

(C)

STRANDEDNESS: double

(D) TOPOLOGY:

linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:19:

GT 2

(2) INFORMATION FOR SEQ ID NO:20:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH:

2

(B) TYPE:

nucleic acid

(C) ST

STRANDEDNESS: double

(D) TOPOLOGY:

linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:20:

TC 2

(2) INFORMATION FOR SEQ ID NO:21:

(i) SEQUENCE CHARACTERISTICS:

(A)

) LENGTH:

3

(B) TYPE:

TYPE: nucleic acid
STRANDEDNESS: double

(C) STRANDEDNI
(D) TOPOLOGY:

linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:21:

CGG 3

(2) INFORMATION FOR SEQ ID NO:22:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH:

3

(B) TYPE:

nucleic acid

(C) STRANDEDNESS: double



71

TOPOLOGY: linear (D) SEQUENCE DESCRIPTION: SEQ ID NO:22: (xi) GGT 3 INFORMATION FOR SEQ ID NO:23: (2) SEQUENCE CHARACTERISTICS: (A) LENGTH: nucleic acid (B) TYPE: STRANDEDNESS: double (C) (D) TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO:23: (xi) ATTA 4 INFORMATION FOR SEQ ID NO:24: (2) SEQUENCE CHARACTERISTICS: (i) LENGTH: (A) nucleic acid (B) TYPE: STRANDEDNESS: double (C) linear (D) TOPOLOGY: SEQUENCE DESCRIPTION: SEQ ID NO:24: (xi) CTGG 4 INFORMATION FOR SEQ ID NO:25: (2) SEQUENCE CHARACTERISTICS: (i) LENGTH: (A) nucleic acid TYPE: (B) STRANDEDNESS: double (C) (D) TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO:25: (xi) GTTATGTCTT ATCCCACGGT ATTTAATATG GTTCATTAGG ATGTTTATTT 50 CTTGATTTTG CATATGAGTA TATTACCCCC CCCTCAAAAA AATAAATTAA 100 TTAAAATGAT GGCTTATATA AAATAAAATT TAAAGCAAGG AATCTCAATG 200 GATGTTAAAC AAAATGAGAT TTTGTGAAAG CAATAAATTA TTGACTTCGT TTTAGATTIG TTTAGCTATA ATGTTATACA TTCAAATGAC TGAACATCCT 250 259 GTAATTAAA INFORMATION FOR SEQ ID NO:26: (2) SEQUENCE CHARACTERISTICS: 256 LENGTH: nucleic acid (B) TYPE: STRANDEDNESS: double (C) linear (D) TOPOLOGY: SEQUENCE DESCRIPTION: SEQ ID NO:26: (xi) TTGTTATGTC TTATCCCACG GTATTTAATA TAGTTCATTT GGATGTTCAT TTCTTTATTT TGCATATGAG TATATTACCC CTTCAAAAAA TAAATTAATT 100 AAAACGATTG CTTATATAAA ACAAAATTTA AAGCAAGGAA TCTCAATGGA 150 TGTTAAACAA AATGAGATTT AGTGAAAACA ATAAATTATT CACTTCGTTT 200

(2) INFORMATION FOR SEQ ID NO:27:

AATTTA

SEQUENCE CHARACTERISTICS:

LENGTH: (A)

nucleic acid (B) TYPE:

TAGATTTGTT TAGCTATAAT GTTATACATT CAAATGACTG AACATCCTGT 250

STRANDEDNESS: double (C) TOPOLOGY: linear (D)

SEQUENCE DESCRIPTION: SEQ ID NO:27: TTTGTTATGT CTTATCCCAC GGTATTTAAT ATAGTTCATT TGGATGTTCA

256



			CCITCINNUL III.	100
			Publiculations in the transfer	150
			Milianii ionollol	200
TTAGATTTGT T	TAGCTATAA	TGTTATACAT	10.22.10.101	250
TAATTAA				257
(2) INFO	RMATION FOR	R SEQ ID NO	:28:	
(i)	SEQUENC	E CHARACTER	RISTICS:	
	(A)	LENGTH:	263	
	(B)	TYPE:	nucleic acid	
	(C)	STRANDEDNE	SS: double	
	(D)	TOPOLOGY:	linear	
(xi)	SEQUENC	E DESCRIPT	ON: SEQ ID NO:28:	
TCTATGTTCT	TATCNCCACG	GTXTTTAATA	TGGTTCATTA GGATGTTTAT	50
TTCTTGATTT	TGCATATGAG	TATATTACCC	CCCCCCTCA AAAAAATAAA	100
TTAATTAAAA	TGATGGCTTA	XATAAAXTAT	AATTTAAAGC AAGGAXTCTC	150
			AAXGCXATXX ATTATTGXCT	200
TCGTTGTAXA	TTTGCTXAGC	TATAATGTTA	TXCATTCAAA TGACTGAACA	250
TCCTGTXXTT				263
(2) INFO	ORMATION FO	R SEQ ID NO):29:	
(i)	SEQUENC	CE CHARACTE	RISTICS:	
		LENGTH:	264	
	(B)	TYPE:	nucleic acid	
			ESS: double	
	(D)	TOPOLOGY:	linear	
(xi) SEQUEN	CE DESCRIPT	ION: SEQ ID NO:29:	
XXXXXXXXX	XTXXXCCCAC	GGTATTTAAT	ATXGTTCATT XGGATGTTXA	50
TTTCTTXATT	TTGCATATGA	GTATATTAXX	XXXXCCCCTX XAAAAAATAA	100
			AAATTTAAAG CAAGGAATCT	150
			GAAAXCAATA AATTATTXAC	200
			ATACATTCAA ATGACTGAAC	250
ATCCTGTAAT				264
(2) INF	ORMATION FO	OR SEQ ID N	0:30:	
(i)	SEQUEN	CE CHARACTE	ERISTICS:	
	(A)	LENGTH:	181	
	(B)	TYPE:	nucleic acid	
	(C)	STRANDEDN	ESS: double	
	(D)	TOPOLOGY:	linear	
(xi	i) SEQUEN	CE DESCRIP	rion: SEQ ID NO:30:	
TTTXCCCGGA	AAAAAATAGO	AAAGGGGGG	G GGGCTAATCG GCAGGGAAGG	50
			CAXAATTTCC AGGTCAGACG	100
			C GGXXGGGGTG TCGTACCACA	150
		G AGATTAAGT		181
(2) IN	EORMATION F	OR SEQ ID N	10:31:	
	•	TOR CUADACT		

LENGTH: 181 (A)

nucleic acid (B) TYPE:

STRANDEDNESS: double (C)

TOPOLOGY: linear (D)

SEQUENCE DESCRIPTION: SEQ ID NO:31:

TXTTXXXCGG AAAAAATXG AAAGGGGGGG GGGCTAATCG GCAGGGAAGG 50 CCGCCCCGGA TAGCGGCCGG CAGAAGGAAT CAGAATTTCC AGGTCAGACG 100 GGCTGCAAGT TGCAGACCGT TAAAATCATC GGTTGGGGTG TCGTACCACA 150 CTTTACCTGC CGTCAGCCCG AGATTAAGTT G





(2)	INFORMA	TION FOR	SEQ ID NO:32:			
(i) SEQUENCE CHARACTERISTICS:						
		(A)	LENGTH:	177		
		(B)	TYPE:	nucleic acid		
		(C)	STRANDEDNESS:	double		
		(D)	TOPOLOGY:	linear		
	(xi)	SEQUENC	E DESCRIPTION:	SEQ ID NO:32:		
TTTXCCG	GAA AAA	AATXGAA	AGGGGGGGC TAAT	CGGCAG GGAAGGCCGC	50	
				CCAGGT CAGATGGGCT		
				TGTCGT ACCACACTTT		
		CCGAGAT			177	
(2)	INFORMA	TION FO	R SEQ ID NO:33:			
	(i) SEQUENCE CHARACTERISTICS:					
		(A)	LENGTH:	182		
		(B)	TYPE:	nucleic acid		
			STRANDEDNESS:	double		
		•	TOPOLOGY:	linear		
	(xi)		E DESCRIPTION:	SEQ ID NO:33:		
тхтхссс		-		ATCGGC AGGGAAGGCC	50	
				TTTCCAG GTCAGACGGG		
				GGTGTC GTACCACACT		
			ATTAAAGTTT GG		182	
TIACCIC	ecco ica	GCCCOM				
(2)	TNFORM	ATION FO	R SEQ ID NO:34:			
(2)	(i)		CE CHARACTERIST			
	(1)	·-	LENGTH:	175		
				nucleic acid		
		•	STRANDEDNESS:			
		, - ,		linear		
	()		CE DESCRIPTION:	•		
VOCCE N	(XÌ)				50	
XCGGAAAAAA ATGAAAGGGG GGGGGCTAAT CGGCAGGGAA GGCCGCCCCG GATAGCGGCC GGCAGAAGGA ATCAGAATTT CCAGGTCAGA TGGGCTGCAA						
				CGTACCA CACTTTACCT	100 150	
		GATAAAG		COIRCEA CACITIACCI	175	
GCCGIC	AGCC CGA	GATAAAG	11100			
(2)	TNEODW	ATION FO	R SEQ ID NO:35:			
(2)	(i)		CE CHARACTERIST			
	(1)	_	LENGTH:	174		
		(B)	TYPE:	nucleic acid		
		(C)	STRANDEDNESS:			
		(D)	TOPOLOGY:	linear		
	(xi)		CE DESCRIPTION:			
CCCNAN		_		AGGGAAG GCCGCCCGG	50	
				GTCAGAT GGGCTGCAAG		
				GTACCAC ACTTTACCTG		
		GATAAAGT		orneene nerrineere	174	
CCGICA	GCCG_GA	JA I AAAG I	1100	÷		
(2)	THEODN	NTTON E	OR SEQ ID NO:36	•		
(2)	(i)		CE CHARACTERIST			
	(1)	(A)	LENGTH:	182		
		(B)	TYPE:	nucleic acid		
		(C)	STRANDEDNESS:			
				linear		
	(xi)		ICE DESCRIPTION:			
хххххх		-		GCTAATCG GCAGGGAAGG	50	
				GAATTTCC AGGTCAGAXG		

GGCTGCAAGT TGCAGACCGT TAXAATCATC GGTTGGGGTG TCGTACCACA 150

nucleic acid



CTTTACCTGC CGTCAGCCCG AGATXAAGTT XG

(B)

- INFORMATION FOR SEQ ID NO:37:
 - SEQUENCE CHARACTERISTICS: (i)

(A) LENGTH:

TYPE:

(C) STRANDEDNESS: single

linear (D) TOPOLOGY:

SEQUENCE DESCRIPTION: SEQ ID NO:37: (xi)

GGGGGGGG 8

- INFORMATION FOR SEQ ID NO:38: (2)
 - SEQUENCE CHARACTERISTICS:

(A) LENGTH:

(B) TYPE: nucleic acid

(C)

STRANDEDNESS: single

(D) TOPOLOGY:

SEQUENCE DESCRIPTION: SEQ ID NO:38: (xi) GGGGGGGG 9

INFORMATION FOR SEQ ID NO:39: (2)

SEQUENCE CHARACTERISTICS: (i)

> (A) LENGTH:

TYPE: (B)

nucleic acid

(C)

STRANDEDNESS: single

(D) TOPOLOGY: linear

SEQUENCE DESCRIPTION: SEQ ID NO:39: (xi)

GGGGGGGGGG 10

- INFORMATION FOR SEQ ID NO:40: (2)
 - SEQUENCE CHARACTERISTICS:

LENGTH: (A)

(B) TYPE: nucleic acid

(C)

STRANDEDNESS: single

linear (D) TOPOLOGY:

SEQUENCE DESCRIPTION: SEQ ID NO:40: (xi)

TAAA 4

- INFORMATION FOR SEQ ID NO:41: (2)
 - SEQUENCE CHARACTERISTICS: (i)

(A)

LENGTH:

nucleic acid (B) TYPE:

(C)

STRANDEDNESS: single

(D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:41:

TTAA 4

- (2) INFORMATION FOR SEQ ID NO:42:
 - SEQUENCE CHARACTERISTICS: (i)

LENGTH:

(A)

(B) TYPE:

nucleic acid

(C) (D)

STRANDEDNESS: single TOPOLOGY:

linear

(xi)

SEQUENCE DESCRIPTION: SEQ ID NO:42:

TTAAA 5